

Fig. 1



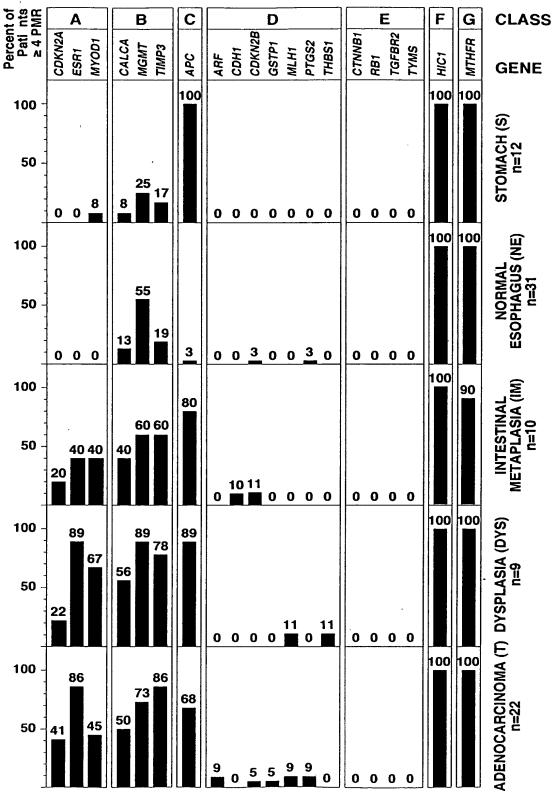
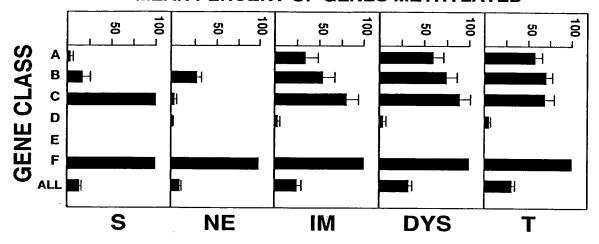


Fig. 2



MEAN PERCENT OF GENES METHYLATED



TISSUE COMPARISON

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<i>p</i> -values	NE \Rightarrow IM	IM → DYS	IM → T	DYS → T
Α	0.0006	0.031	0.015	0.87
В	0.027	0.13	0.15	0.71
С	< 0.0001	0.55	0.34	0.11
D	0.46	0.92	0.40	0.48
E		_		_
F		_		_
ALL	< 0.0001	0.044	0.042	0.70

Fig. 3



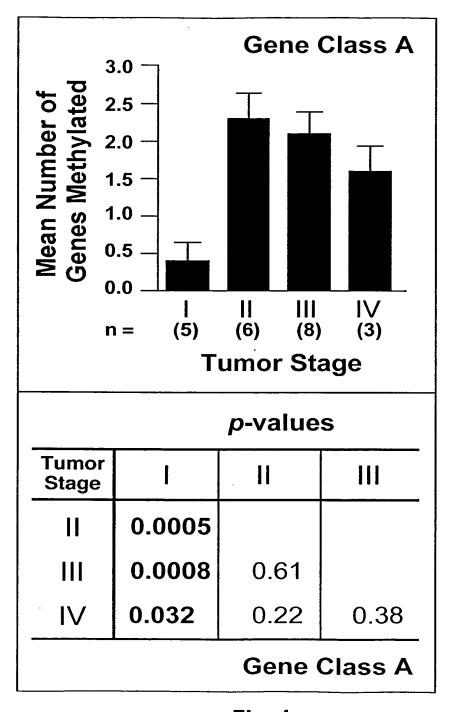
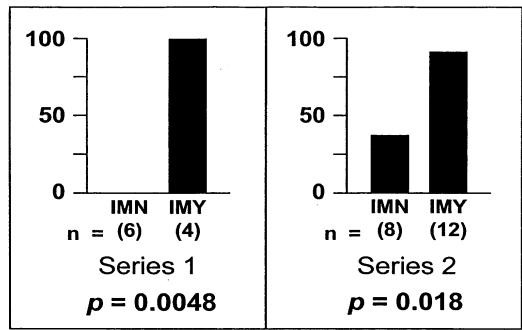


Fig. 4





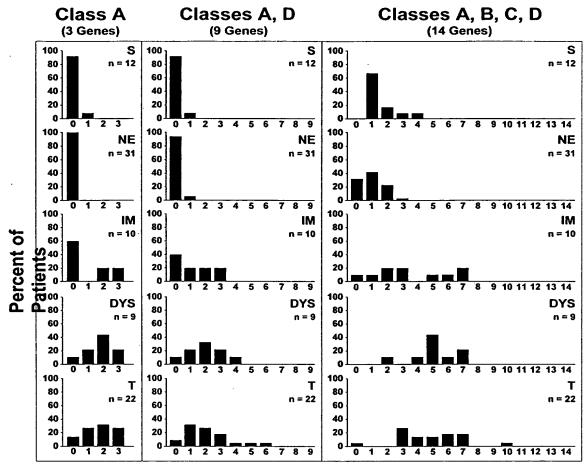


INTESTINAL METAPLASIA (IM)

(IMY = Associated Dysplasia) (IMN = No Associated Dysplasia)

Fig. 5





Number of Methylated CpG Islands

Fig. 6